

FIGURE 2

Best fit versus local and global alignment

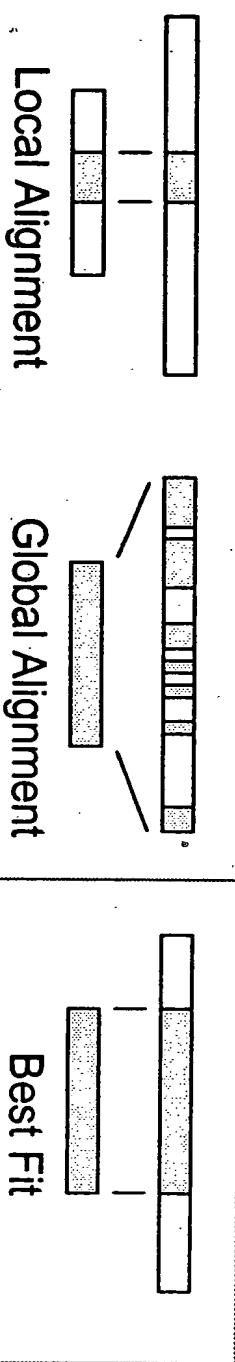


Fig. 3A

Fig. 3B

Fig. 3C

Identity → Gap → Mismatch →

1	MTPGGRSLPPSV	DPHPLLLIGLLPMPLEAQGLRGIGLTPSAAQPAHQ	49
1	MTPGGRLLYLLRWCSTPPPLLLIGLLAIPLEAQGLRGIGLTPSAAQPAHQ	50	
50	LPTPFTRGTLKPAAHLVGDPSPNRTTLRANTDRAFLPTAFSLSNNSLV	99	
51	LPTPFTRGTLKPAAHLVGDPSTQDSLWRANTDRAFLRHHGFSLSNNSLV	100	
100	PTSGLYFVYSQWFSGRGCFPRATPTPLYLAHEVQLFSPQYFHVPLSA	149	
101	PTSGLYFVYSQWFSGRGCFPRATPTPLYLAHEVQLFSPQYFHVPLSA	150	
150	QKSVCQGPQGRWRSVYQGAVFLTRGDQLSTHTDGISHLLSPSSVFFG	199	
151	QKSVCQGPQGPWRSVYQGAVFLTRGDQLSTHTDGISHLLSPSSVFFG	200	
200	AFAL	203	
201	AFAL	204	

FIG. 4